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=> s detect? (3a) (polymorphism? or variant? or mutation?)
4 FILES SEARCHED...

L1 69865 DETECT? (3A) (POLYMORPHISM? OR VARIANT? OR MUTATION?)

=> s 11 and modified base

L2 958 L1 AND MODIFIED BASE

=> s 12 and 90%

L3 859 L2 AND 90%

=> s 13 and 90% (3a)points (4a) occurrence L4 4 L3 AND 90% (3A) POINTS (4A) OCCURRENCE

=> d 14 bib abs 1-4

L4 ANSWER 1 OF 4 USPATFULL on STN

AN 2003:237723 USPATFULL

TI Fluorescence-based genotyping

IN Wolfe, Jia Liu, Winchester, MA, UNITED STATES
Kawate, Tomohiko, Cambridge, MA, UNITED STATES
Allerson, Charles R., Carlsbad, CA, UNITED STATES
Stanton, Vincent P., JR., Belmont, MA, UNITED STATES

PI US 2003165880 A1 20030904

AI US 2002-107748 A1 20020326 (10)

RLI Continuation-in-part of Ser. No. US 1999-394467, filed on 10 Sep 1999, GRANTED, Pat. No. US 6566059

PRAI US 1998-102724P 19981001 (60)

DT Utility

FS APPLICATION

LREP BINGHAM, MCCUTCHEN LLP, THREE EMBARCADERO, SUITE 1800, SAN FRANCISCO, CA, 94111-4067

CLMN Number of Claims: 26 ECL Exemplary Claim: 1

DRWN 13 Drawing Page(s)

LN.CNT 1356

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention relates to a method for genotyping a diploid

organism to **detect** single nucleotide **polymorphisms** (SNPs) using modified nucleotides or nucleotide residues substituted with fluorescent groups.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

```
ANSWER 2 OF 4 USPATFULL on STN
ΑN
       2003:228236 USPATFULL
TI
       Base-modified nucleotides and cleavage of polynucleotides incorporating
       them
       Stanton, Jr., Vincent P., Belmont, MA, United States
IN
       Wolfe, Jia Liu, Winchester, MA, United States
       Kawate, Tomohiko, Cambridge, MA, United States
       Allerson, Charles, Cambridge, MA, United States
       Verdine, Gregory L., Cambridge, MA, United States
PA
       Variagenics, Inc., Cambridge, MA, United States (U.S. corporation)
PI
       US 6610492
                               20030826
                          В1
       US 2002-43511
ΑI
                               20020108 (10)
RLI
       Continuation-in-part of Ser. No. US 1999-394467, filed on 10 Sep 1999
PRAI
       US 1998-102724P
                          19981001 (60)
DТ
       Utility
FS
       GRANTED
EXNAM
       Primary Examiner: Riley, Jezia
LREP
       Rose, Esq., Bernard F., Bingham McCutchen LLP
CLMN
       Number of Claims: 15
ECL
       Exemplary Claim: 1
DRWN
       12 Drawing Figure(s); 12 Drawing Page(s)
LN.CNT 1820
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
       The present invention is directed to novel base-modified nucleotides and
       methods for their use in the preparation and cleavage of modified
       polynucleotides.
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
L4
     ANSWER 3 OF 4 USPATFULL on STN
       2003:136911 USPATFULL
ΑN
TΙ
       Method for analyzing polynucleotides
IN
       Stanton, Jr., Vincent P., Belmont, MA, United States
       Wolfe, Jia Liu, Winchester, MA, United States
       Verdine, Gregory L., Cambridge, MA, United States
PΑ
       Variagenics, Inc., Cambridge, MA, United States (U.S. corporation)
PΤ
       US 6566059
                               20030520
                          B1
       US 1999-394467
ΑI
                               19990910 (9)
       US 1998-102724P
PRAI
                           19981001 (60)
       US 1999-149533P
                           19990817 (60)
DT
       Utility
FS
       GRANTED
      Primary Examiner: Riley, Jezia
EXNAM
LREP
       Rose, Bernard F., Bingham McCutchen LLP
СТМИ
       Number of Claims: 69
ECL
       Exemplary Claim: 1
DRWN
       52 Drawing Figure(s); 39 Drawing Page(s)
LN.CNT 5806
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
       The present invention relates to methods for the analysis of
AB
       polynucleotides including detection of variance in nucleotide sequence
       without the need for full sequence determination, full sequence
       determination of a polynucleotide, genotyping of DNA and labeling a
       polynucleotide fragment during the process of cleaving it into
       fragments.
```

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 4 OF 4 USPATFULL on STN T.4 AN2002:217059 USPATFULL Method for analyzing polynucleotides ΤI Stanton, Jr., Vincent P., 32 Royal Rd., Belmont, MA, United States IN Wolfe, Jia Liu, 28 Hollywood Rd., Winchester, MA, United States 01890 Kawate, Tomohiko, 31 Portsmouth St., 1R, Cambridge, MA, United States Verdine, Gregory L., 7 Bennington Rd., Lexington, MA, United States 02421 US 6440705 В1 20020827 ΡI US 1999-394457 19990910 (9) AΙ 19990817 (60) PRAI US 1999-149533P US 1998-102724P 19981001 (60) DT Utility FS GRANTED EXNAM Primary Examiner: Siew, Jeffrey LREP Lyon & Lyon LLP Number of Claims: 32 CLMN ECLExemplary Claim: 1 52 Drawing Figure(s); 39 Drawing Page(s) DRWN LN.CNT 5788 CAS INDEXING IS AVAILABLE FOR THIS PATENT. The present invention relates to methods for the analysis of polynucleotides including detection of variance in nucleotide sequence without the need for full sequence determination, full sequence determination of a polynucleotide, genotyping of DNA and labeling a polynucleotide fragment during the process of cleaving it into fragments.

CAS INDEXING IS AVAILABLE FOR THI